

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/719,587 A
Source: IFW16
Date Processed by STIC: 08/17/2006

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 08/17/2006

PATENT APPLICATION: US/10/719,587A

TIME: 11:06:15

Input Set : A:\55862-CIP(46342)-Amended.txt

Output Set: N:\CRF4\08172006\J719587A.raw

3 <110> APPLICANT: Takeda Chemical Industries, Ltd.
 5 <120> TITLE OF INVENTION: Novel G protein-coupled receptor protein, its DNA and ligand
 thereof
 7 <130> FILE REFERENCE: 2568USOP-CIP
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/719,587A
 C--> 9 <141> CURRENT FILING DATE: 2003-11-21
 9 <150> PRIOR APPLICATION NUMBER: US 09/831,758
 10 <151> PRIOR FILING DATE: 2001-05-11
 12 <150> PRIOR APPLICATION NUMBER: PCT/JP99/06283
 13 <151> PRIOR FILING DATE: 1999-11-11
 15 <150> PRIOR APPLICATION NUMBER: JP 10-323759
 16 <151> PRIOR FILING DATE: 1998-11-13
 18 <150> PRIOR APPLICATION NUMBER: JP 11-060030
 19 <151> PRIOR FILING DATE: 1999-03-08
 21 <150> PRIOR APPLICATION NUMBER: JP 11-106812
 22 <151> PRIOR FILING DATE: 1999-04-14
 24 <150> PRIOR APPLICATION NUMBER: JP 11-166672
 25 <151> PRIOR FILING DATE: 1999-06-14
 27 <150> PRIOR APPLICATION NUMBER: JP 11-221640
 28 <151> PRIOR FILING DATE: 1999-08-04
 30 <150> PRIOR APPLICATION NUMBER: JP 11-259818
 31 <151> PRIOR FILING DATE: 1999-09-14
 33 <160> NUMBER OF SEQ ID NOS: 66
 35 <210> SEQ ID NO: 1
 36 <211> LENGTH: 180
 37 <212> TYPE: PRT
 38 <213> ORGANISM: Human
 40 <400> SEQUENCE: 1
 41 Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr
 42 1 5 10 15
 43 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Met
 44 20 25 30
 45 Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
 46 35 40 45
 47 Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
 48 50 55 60
 49 Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
 50 65 70 75 80
 51 Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
 52 85 90 95
 53 Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
 54 100 105 110
 55 Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
 56 115 120 125

CP9-6)

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57 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
58      130                      135                      140
59 Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
60 145                      150                      155                      160
61 Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
62                      165                      170                      175
63 Lys Gln Ser Arg
64      180
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 540
68 <212> TYPE: DNA
69 <213> ORGANISM: Human
71 <400> SEQUENCE: 2
72 atggaaatta tttcatcaaa actattcatt ttattgactt tagccacttc aagcttggtta 60
73 acatcaaaca ttttttgtgc agatgaatta gtgatgtcca atcttcacag caaagaaaaat 120
74 tatgacaaat attctgagcc tagaggatac caaaaagggg aaagaagcct caattttgag 180
75 gaattaaaag attgggggacc aaaaaatgtt attaagatga gtacacctgc agtcaataaa 240
76 atgccacact ccttcgccaa cttgccattg agatttgga ggaacgttca agaagaaaga 300
77 agtgctggag caacagccaa cctgcctctg agatctgga agaaatatgga ggtgagcttc 360
78 gtgagacgtg ttcctaacct gcccacaaagg tttgggagaa caacaacagc caaaagtgtc 420
79 tgcaggatgc tgagtattt gtgtcaagga tccatgcatt caccatgtgc caatgactta 480
80 ttttactcca tgacctgcca gcaccaagaa atccagaatc ccgatcaaaa acagtcaagg 540
82 <210> SEQ ID NO: 3
83 <211> LENGTH: 27
84 <212> TYPE: DNA
85 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: primer
90 <400> SEQUENCE: 3
91 gggctgcaca tagagactta atttttag 27
93 <210> SEQ ID NO: 4
94 <211> LENGTH: 27
95 <212> TYPE: DNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: primer
101 <400> SEQUENCE: 4
102 ctagaccacc tctatataac tgcccat 27
104 <210> SEQ ID NO: 5
105 <211> LENGTH: 30
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: primer
112 <400> SEQUENCE: 5
113 gcacatagag acttaatttt agatttagac 30
115 <210> SEQ ID NO: 6
116 <211> LENGTH: 27
117 <212> TYPE: DNA

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118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <223> OTHER INFORMATION: primer
123 <400> SEQUENCE: 6
124 catgcacttt gactggtttc caggtat 27
126 <210> SEQ ID NO: 7
127 <211> LENGTH: 27
128 <212> TYPE: DNA
129 <213> ORGANISM: Artificial Sequence
131 <220> FEATURE:
132 <223> OTHER INFORMATION: primer
134 <400> SEQUENCE: 7
135 cagctttagg gacaggctcc aggtttc 27
137 <210> SEQ ID NO: 8
138 <211> LENGTH: 196
139 <212> TYPE: PRT
140 <213> ORGANISM: Human
142 <400> SEQUENCE: 8
143 Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr
144 1 5 10 15
145 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Met
146 20 25 30
147 Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
148 35 40 45
149 Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
150 50 55 60
151 Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
152 65 70 75 80
153 Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
154 85 90 95
155 Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
156 100 105 110
157 Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
158 115 120 125
159 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
160 130 135 140
161 Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
162 145 150 155 160
163 Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
164 165 170 175
165 Lys Gln Ser Arg Arg Leu Leu Phe Lys Lys Ile Asp Asp Ala Glu Leu
166 180 185 190
167 Lys Gln Glu Lys
168 195
170 <210> SEQ ID NO: 9
171 <211> LENGTH: 588
172 <212> TYPE: DNA
173 <213> ORGANISM: Human
175 <400> SEQUENCE: 9

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```

176 atggaaatta tttcatcaaa actattcatt ttattgactt tagccacttc aagcttggtta 60
177 acatcaaaca ttttttgtgc agatgaatta gtgatgtcca atcttcacag caaagaaaat 120
178 tatgacaaat attctgagcc tagaggatac ccaaaagggg aaagaagcct caattttgag 180
179 gaattaaaag attggggacc aaaaaatggt attaagatga gtacacctgc agtcaataaa 240
180 atgccacact ccttcgccaa cttgccattg agatttggga ggaacgttca agaagaaaga 300
181 agtgctggag caacagccaa cctgcctctg agatctggaa gaaatatgga ggtgagcctc 360
182 gtgagacgtg ttcctaacct gccccaaagg tttgggagaa caacaacagc caaaagtgtc 420
183 tgcaggatgc tgagtgtatt gtgtcaagga tccatgcatt caccatgtgc caatgactta 480
184 ttttactcca tgacctgcca gcaccaagaa atccagaatc ccgatcaaaa acagtcaagg 540
185 agactgctat tcaagaaaat agatgatgca gaattgaaac aagaaaaa 588
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 27
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial Sequence
192 <220> FEATURE:
193 <223> OTHER INFORMATION: primer
195 <400> SEQUENCE: 10
196 gcctagagga gatctaggct gggagga 27
198 <210> SEQ ID NO: 11
199 <211> LENGTH: 27
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: primer
206 <400> SEQUENCE: 11
207 gggaggaaca tggaagaaga aaggagc 27
209 <210> SEQ ID NO: 12
210 <211> LENGTH: 27
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: primer
217 <400> SEQUENCE: 12
218 gatggtgaat gcatggactg ctggagc 27
220 <210> SEQ ID NO: 13
221 <211> LENGTH: 27
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: primer
228 <400> SEQUENCE: 13
229 ttcctcccaa atctcagtg caggttg 27
231 <210> SEQ ID NO: 14
232 <211> LENGTH: 196
233 <212> TYPE: PRT
234 <213> ORGANISM: Bovine
236 <400> SEQUENCE: 14
237 Met Glu Ile Ile Ser Leu Lys Arg Phe Ile Leu Leu Met Leu Ala Thr
238 1 5 10 15

```

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```

239 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Thr Asp Glu Ser Arg Met
240          20          25          30
241 Pro Asn Leu Tyr Ser Lys Lys Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
242          35          40          45
243 Gly Asp Leu Gly Trp Glu Lys Glu Arg Ser Leu Thr Phe Glu Glu Val
244          50          55          60
245 Lys Asp Trp Ala Pro Lys Ile Lys Met Asn Lys Pro Val Val Asn Lys
246 65          70          75          80
247 Met Pro Pro Ser Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Met
248          85          90          95
249 Glu Glu Glu Arg Ser Thr Arg Ala Met Ala His Leu Pro Leu Arg Leu
250          100         105         110
251 Gly Lys Asn Arg Glu Asp Ser Leu Ser Arg Trp Val Pro Asn Leu Pro
252          115         120         125
253 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Ile Thr Lys Thr Leu
254          130         135         140
255 Ser Asn Leu Leu Gln Gln Ser Met His Ser Pro Ser Thr Asn Gly Leu
256 145         150         155         160
257 Leu Tyr Ser Met Ala Cys Gln Pro Gln Glu Ile Gln Asn Pro Gly Gln
258          165         170         175
259 Lys Asn Leu Arg Arg Arg Gly Phe Gln Lys Ile Asp Asp Ala Glu Leu
260          180         185         190
261 Lys Gln Glu Lys
262          195

```

264 <210> SEQ ID NO: 15

265 <211> LENGTH: 588

266 <212> TYPE: DNA

267 <213> ORGANISM: Bovine

269 <400> SEQUENCE: 15

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270 atggaaatta ttctattaaa acgattcatt ttattgatgt tagccacttc aagcttggtta 60
271 acatcaaaca tcttctgcac agacgaatca aggatgccca atctttacag caaaaagaat 120
272 tatgacaaat attccgagcc tagaggagat ctaggctggg agaaagaaag aagtcttact 180
273 tttgaagaag taaaagattg ggctccaaaa attaagatga ataaacctgt agtcaacaaa 240
274 atgccacctt ctgcagccaa cctgccactg agatttgagg ggaacatgga agaagaaagg 300
275 agcactaggg cgatggccca cctgcctctg agactcggaa aaaatagaga ggacagcctc 360
276 tccagatggg tcccaaattc gcccagagg tttggaagaa caacaacagc caaaagcatt 420
277 accaagaccc tgagtaattt gctccagcag tccatgcatt caccatctac caatgggcta 480
278 ctctactcca tggcctgcca gcccgaagaa atccagaatc ctggtcaaaa gaacctaaag 540
279 agacggggat tccagaaaat agatgatgca gaattgaaac aagaaaaaa 588

```

281 <210> SEQ ID NO: 16

282 <211> LENGTH: 27

283 <212> TYPE: DNA

284 <213> ORGANISM: Artificial Sequence

286 <220> FEATURE:

287 <223> OTHER INFORMATION: primer

289 <400> SEQUENCE: 16

290 ccctggggct tcttctgtct tctatgt

27

292 <210> SEQ ID NO: 17

293 <211> LENGTH: 26

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/17/2006
PATENT APPLICATION: US/10/719,587A TIME: 11:06:16

Input Set : A:\55862-CIP(46342)-Amended.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:20; N Pos. 3,9
Seq#:21; N Pos. 3,9,12
Seq#:22; N Pos. 3,6,9
Seq#:23; N Pos. 3,6,9,12
Seq#:24; N Pos. 3,6,9
Seq#:25; N Pos. 3,6,9,12
Seq#:63; Xaa Pos. 10

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:64,65,66

VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:456 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:1145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0